

LAP13 Rec'd PCT/PTO 14 FEB 2007

UNITED STATES PATENT AND TRADEMARK OFFICE

Applicant: Fukatsu et al. Examiner:
Serial No.: 10/580906 Group Art: unknown
Filed: May 26, 2006 Docket: 20039.0005USWO

Title: RECEPTOR FUNCTION REGULATING AGENT

CERTIFICATE UNDER 37 CFR 1.10

Express Mail mailing label number: EV 802672995 US

Date of Deposit: February 14, 2007

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Sir:

The following papers are transmitted herewith:

- ☒ Transmittal Sheet in duplicate containing Certificate of Mailing 1.10
- ☒ Sequence listing diskette (computer readable); sequence listing (paper copy)
- ☒ Communication: Submission of Computer Readable Sequence Listing
- ☒ Copy of the Notification to Comply with Requirements mailed December 14, 2006
- ☒ Return Postcard

Please charge any additional fees or credit overpayment to Deposit Account No. 50-3478. A duplicate of this sheet is enclosed.

Hamre, Schumann, Mueller & Larson, P.C.

P.O. Box 2902 Minneapolis, MN 55402

612.455-3800

By: 

Name: Douglas P. Mueller

Reg. No.: 30,300

Initials: DPM:rkw



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U.S. APPLICATION NUMBER NO. 10/580,906	FIRST NAMED APPLICANT Kohji Fukatsu	ATTY. DOCKET NO. 20039.0005USWO
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INTERNATIONAL APPLICATION NO. PCT/JP04/17996	
LA. FILING DATE 11/26/2004	PRIORITY DATE 11/26/2003

52835
HAMRE, SCHUMANN, MUELLER & LARSON, P.C.
P.O. BOX 2902
MINNEAPOLIS, MN 55402-0902

CONFIRMATION NO. 7379
371 FORMALITIES LETTER



Date Mailed: 12/14/2006

NOTIFICATION TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant is given **TWO MONTHS FROM THE DATE OF THIS NOTICE** within which to file the items indicated below to avoid abandonment. Extensions of time may be obtained under the provisions of 37 CFR 1.136(a).

- A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

Applicant is cautioned that correction of the above items may cause the specification and drawings page count to exceed 100 pages. If the specification and drawings exceed 100 pages, applicant will need to submit the required application size fee.

For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:

- For Rules Interpretation, call (571) 272-0951
- For Patentin Software Program Help, call Patent EBC at 1-866-217-9197 or directly at 703-305-3028 / 703-308-6845 between the hours of 6 a.m. and 12 midnight, Monday through Friday, EST.
- Send e-mail correspondence for Patentin Software Program Help @ ebc@uspto.gov

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

Registered users of EFS-Web may alternatively submit their reply to this notice via EFS-Web.
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For more information about EFS-Web please call the USPTO Electronic Business Center at 1-866-217-9197 or visit our website at <http://www.uspto.gov/ebc>.

Seq. Digtette: 2/14/2007

10

If you are not using EFS-Web to submit your reply, you must include a copy of this notice.

BARBARA A CAMPBELL

Telephone: (703) 308-9140 EXT 217

PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
10/580,906	PCT/JP04/17996	20039.0005USWO

FORM PCT/DO/EO/922 (371 Formalities Notice)

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/580,906
Source: 194P
Date Processed by STIC: 6/8/06

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

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Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/efc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWP

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/580,906

DATE: 06/08/2006
TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

3 <110> APPLICANT: FUKATSU et al.
5 <120> TITLE OF INVENTION: RECEPTOR FUNCTION REGULATING AGENT
7 <130> FILE REFERENCE: 20039.0005USWO
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/580,906
10 <141> CURRENT FILING DATE: 2006-05-26
12 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/017996
13 <151> PRIOR FILING DATE: 2004-11-26
15 <150> PRIOR APPLICATION NUMBER: JP 2003-394848
16 <151> PRIOR FILING DATE: 2003-11-26
18 <160> NUMBER OF SEQ ID NOS: 20
20 <170> SOFTWARE: PatentIn Version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 361
24 <212> TYPE: PRT
25 <213> ORGANISM: Human
W--> 26 <400> SEQUENCE: 1
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29 5 10 15
30 Leu Glu Gln Ala Asn Arg Thr Arg Phe Pro Phe Phe Ser Asp Val Lys
31 20 25 30
32 Gly Asp His Arg Leu Val Leu Ala Ala Val Glu Thr Thr Val Leu Val
33 35 40 45
34 Leu Ile Phe Ala Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
35 50 55 60
36 Val Ala Arg Arg Arg Arg Gly Ala Thr Ala Cys Leu Val Leu Asn
37 65 70 75 80
38 Leu Phe Cys Ala Asp Leu Leu Phe Ile Ser Ala Ile Pro Leu Val Leu
39 85 90 95
40 Ala Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Ala Cys His
41 100 105 110
42 Leu Leu Phe Tyr Val Met Thr Leu Ser Gly Ser Val Thr Ile Leu Thr
43 115 120 125
44 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val His Leu Gln
45 130 135 140
46 Arg Gly Val Arg Gly Pro Gly Arg Arg Ala Arg Ala Val Leu Leu Ala
47 145 150 155 160
48 Leu Ile Trp Gly Tyr Ser Ala Val Ala Ala Leu Pro Leu Cys Val Phe
49 165 170 175
50 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Ala Asp Gln Glu Ile Ser
51 180 185 190
52 Ile Cys Thr Leu Ile Trp Pro Thr Ile Pro Gly Glu Ile Ser Trp Asp
53 195 200 205
54 Val Ser Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val

pg 4,6
Does Not Comply
Corrected Diskette Needed

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/580,906

DATE: 06/08/2006
TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

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55      210      215      220
56 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
57 225      230      235      240
58 Leu Thr Val Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
59      245      250      255
60 Gln Gln Asp Phe Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
61      260      265      270
62 Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
63      275      280      285
64 Ile Gln Asn Phe Lys Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
65      290      295      300
66 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
67 305      310      315      320
68 Tyr Asn Met Thr Leu Cys Arg Asn Glu Trp Lys Lys Ile Phe Cys Cys
69      325      330      335
70 Phe Trp Phe Pro Glu Lys Gly Ala Ile Leu Thr Asp Thr Ser Val Lys
71      340      345      350
72 Arg Asn Asp Leu Ser Ile Ile Ser Gly
73      355      360

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74 <210> SEQ ID NO: 2

75 <211> LENGTH: 1083

76 <212> TYPE: DNA

77 <213> ORGANISM: Human

N--> 78 <400> SEQUENCE: 2

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80 aaccgcaccc gctttccctt cttctccgac gtcaagggcg accaccggct ggtgctggcc 120
81 gcggtggaga caaccgtgct ggtgctcatc tttgcagtgt cgctgctggg caacgtgtgc 180
82 gccctggtgc tgggtggcgc cgcagcagc cgcggcgcca ctgacctgct ggtactcaac 240
83 ctcttctgcg cggacctgct cttcatcagc gctatccctc tgggtgctggc cgtgcgctgg 300
84 actgaggcct ggctgctggg ccccggttgc tgcacctgc tcttctacgt gatgacctg 360
85 agcggcagcg tcaccatcct cagctggcc gcggtcagcc tggagcgcat ggtgtgcatc 420
86 gtgcacctgc agcgcggcgt gcggggtcct gggcggcggg cgcgggcagt gctgctggcg 480
87 ctcatctggg gctattcggc ggtcgccgct ctgacctctt gcgtcttctt ccgagtcgct 540
88 ccgcaacggc tccccggcgc cgaccaggaa atttcgattt gcacactgat ttggcccacc 600
89 attcctggag agatctcgtg ggatgtctct tttgttactt tgaacttctt ggtgccagga 660
90 ctggtcattg tgatcagtta ctccaaaatt ttacagatca caaaggcatc aaggaagagg 720
91 ctacaggtaa gcctggccta ctcgagagc caccagatcc gcgtgtccca gcaggacttc 780
92 cggctcttcc gcacctctt cctcctcatg gtctccttct tcatcatgtg gagccccatc 840
93 atcatcacca tctcctcat cctgatccag aacttcaagc aagacctggg catctggccg 900
94 tccctcttct tctgggtggg ggccttcaca tttgctaatt cagccctaaa ccccatcctc 960
95 tacaacatga cactgtgcag gaatgagtgg aagaaaattt tttgctgctt ctggttccca 1020
96 gaaaaggagg ccattttaac agacacatct gtcaaaagaa atgacttgct gattatttct 1080
97 ggc 1083

```

98 <210> SEQ ID NO: 3

99 <211> LENGTH: 361

100 <212> TYPE: PRT

101 <213> ORGANISM: Mouse

N--> 102 <400> SEQUENCE: 3

103 Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser His Thr

RAW SEQUENCE LISTING

DATE: 06/08/2006

PATENT APPLICATION: US/10/580,906

TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

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104          5          10          15
105 Leu Asp Gln Val Asn Arg Thr His Phe Pro Phe Phe Ser Asp Val Lys
106          20          25          30
107 Gly Asp His Arg Leu Val Leu Ser Val Val Glu Thr Thr Val Leu Gly
108          35          40          45
109 Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
110          50          55          60
111 Val Ala Arg Arg Arg Arg Arg Gly Ala Thr Ala Ser Leu Val Leu Asn
112 65          70          75          80
113 Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
114          85          90          95
115 Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
116          100          105          110
117 Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
118          115          120          125
119 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
120          130          135          140
121 Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
122 145          150          155          160
123 Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
124          165          170          175
125 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
126          180          185          190
127 Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
128          195          200          205
129 Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
130          210          215          220
131 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
132 225          230          235          240
133 Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
134          245          250          255
135 Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
136          260          265          270
137 Phe Phe Ile Met Trp Ser Pro Ile Ile Thr Ile Leu Leu Ile Leu
138          275          280          285
139 Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
140          290          295          300
141 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
142 305          310          315          320
143 Tyr Asn Met Ser Leu Phe Arg Asn Glu Trp Arg Lys Ile Phe Cys Cys
144          325          330          335
145 Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Asp Thr Ser Val Arg
146          340          345          350
147 Arg Asn Asp Leu Ser Val Ile Ser Ser
148          355          360
149 <210> SEQ ID NO: 4
150 <211> LENGTH: 1083
151 <212> TYPE: DNA
152 <213> ORGANISM: Mouse

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RAW SEQUENCE LISTING

DATE: 06/08/2006

PATENT APPLICATION: US/10/580,906

TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

W--> 153 <400> SEQUENCE: 4

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154 atgtccctg agtgtgcaca gacgacgggc cctggccct cgcacaccct ggaccaagtc 60
155 aatcgacccc acttcccttt cttctcgat gtcaaggcg accaccggtt ggtgttgagc 120
156 gtcgtggaga ccaccgttct ggggctcatc tttgtcgtct cactgctggg caacgtgtgt 180
157 gctctagtgc tgggtggcgcg ccgtcggcgc cgtggggcga cagccagcct ggtgctcaac 240
158 ctcttctgcg cggatttgct cttcaccage gccatccctc tagtgctcgt cgtgcgctgg 300
159 actgaggcct ggtgttggg gcccgctcgtc tgccacctgc tcttctacgt gatgacaatg 360
160 agcggcagcg tcacgatcct cacactggcc gcggtcagcc tggagcgcac ggtgtgcatc 420
161 gtgcgcctcc ggcgcggctt gagcggcccg gggcggcgga ctcaggcggc actgctggct 480
162 ttcatatggg gttactcggc gctcgccgcg ctgcccctct gcatcttgtt ccgcgtggtc 540
163 ccgcagcgcc tccccggcgg ggaccaggaa attccgattt gcacattgga ttggcccaac 600
164 cgcataggag aaatctcatg ggatgtgttt tttgtgactt tgaacttcct ggtgccggga 660
165 ctggtcattg tgatcagtta ctccaaaatt ttacagatca cgaaagcatc gcggaagagg 720
166 cttacgctga gcttggcata ctctgagagc caccagatcc gagtgtccca acaagactac 780
167 cgactcttcc gcacgctctt cctgctcatg gtttcttct tcatcatgtg gagtcccatc 840
168 atcatcacca tcctctcat cttgatccaa aacttcggc aggacctggt catctggcca 900
169 tcccccttct tctgggtggt ggccttcacg tttgccaact ctgccctaaa cccatactg 960
170 tacaacatgt cgctgttcag gaacgaatgg aggaagattt tttgctgctt cttttttcca 1020
171 gagaagggag ccatttttac agacacgtct gtcaggcgaa atgacttgte tgttatttcc 1080
172 agc 1083

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173 <210> SEQ ID NO: 5

174 <211> LENGTH: 20

175 <212> TYPE: DNA

176 <213> ORGANISM: Artificial Sequence

W--> 177 <220> FEATURE:

W--> 178 <223> OTHER INFORMATION:

W--> 178 <400> SEQUENCE: 5

179 gctgtggcat gcttttaaac 20

180 <210> SEQ ID NO: 6

181 <211> LENGTH: 20

182 <212> TYPE: DNA

183 <213> ORGANISM: Artificial Sequence

W--> 184 <220> FEATURE:

W--> 185 <223> OTHER INFORMATION:

W--> 185 <400> SEQUENCE: 6

186 cgctgtggat gcttatttgc 20

187 <210> SEQ ID NO: 7

188 <211> LENGTH: 30

189 <212> TYPE: DNA

190 <213> ORGANISM: Artificial Sequence

W--> 191 <220> FEATURE:

W--> 192 <223> OTHER INFORMATION:

W--> 192 <400> SEQUENCE: 7

193 agttcatttc cagtaccctc catcagtggc 30

194 <210> SEQ ID NO: 8

195 <211> LENGTH: 361

196 <212> TYPE: PRT

197 <213> ORGANISM: Rat

W--> 198 <400> SEQUENCE: 8

*see p. 6 for env
exploration*

*this env appears in
other sequences too*

RAW SEQUENCE LISTING

DATE: 06/08/2006

PATENT APPLICATION: US/10/580,906

TIME: 10:08:05

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

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199 Met Ser Pro Glu Cys Ala Gln Thr Thr Gly Pro Gly Pro Ser Arg Thr
200           5           10           15
201 Pro Asp Gln Val Asn Arg Thr His Phe Pro Phe Phe Ser Asp Val Lys
202           20           25           30
203 Gly Asp His Arg Leu Val Leu Ser Val Leu Glu Thr Thr Val Leu Gly
204           35           40           45
205 Leu Ile Phe Val Val Ser Leu Leu Gly Asn Val Cys Ala Leu Val Leu
206           50           55           60
207 Val Val Arg Arg Arg Arg Arg Gly Ala Thr Val Ser Leu Val Leu Asn
208           65           70           75           80
209 Leu Phe Cys Ala Asp Leu Leu Phe Thr Ser Ala Ile Pro Leu Val Leu
210           85           90           95
211 Val Val Arg Trp Thr Glu Ala Trp Leu Leu Gly Pro Val Val Cys His
212           100          105          110
213 Leu Leu Phe Tyr Val Met Thr Met Ser Gly Ser Val Thr Ile Leu Thr
214           115          120          125
215 Leu Ala Ala Val Ser Leu Glu Arg Met Val Cys Ile Val Arg Leu Arg
216           130          135          140
217 Arg Gly Leu Ser Gly Pro Gly Arg Arg Thr Gln Ala Ala Leu Leu Ala
218 145           150          155          160
219 Phe Ile Trp Gly Tyr Ser Ala Leu Ala Ala Leu Pro Leu Cys Ile Leu
220           165          170          175
221 Phe Arg Val Val Pro Gln Arg Leu Pro Gly Gly Asp Gln Glu Ile Pro
222           180          185          190
223 Ile Cys Thr Leu Asp Trp Pro Asn Arg Ile Gly Glu Ile Ser Trp Asp
224           195          200          205
225 Val Phe Phe Val Thr Leu Asn Phe Leu Val Pro Gly Leu Val Ile Val
226           210          215          220
227 Ile Ser Tyr Ser Lys Ile Leu Gln Ile Thr Lys Ala Ser Arg Lys Arg
228 225           230          235          240
229 Leu Thr Leu Ser Leu Ala Tyr Ser Glu Ser His Gln Ile Arg Val Ser
230           245          250          255
231 Gln Gln Asp Tyr Arg Leu Phe Arg Thr Leu Phe Leu Leu Met Val Ser
232           260          265          270
233 Phe Phe Ile Met Trp Ser Pro Ile Ile Ile Thr Ile Leu Leu Ile Leu
234           275          280          285
235 Ile Gln Asn Phe Arg Gln Asp Leu Val Ile Trp Pro Ser Leu Phe Phe
236           290          295          300
237 Trp Val Val Ala Phe Thr Phe Ala Asn Ser Ala Leu Asn Pro Ile Leu
238 305           310          315          320
239 Tyr Asn Met Ser Leu Phe Arg Ser Glu Trp Arg Lys Ile Phe Cys Cys
240           325          330          335
241 Phe Phe Phe Pro Glu Lys Gly Ala Ile Phe Thr Glu Thr Ser Ile Arg
242           340          345          350
243 Arg Asn Asp Leu Ser Val Ile Ser Thr
244           355          360
245 <210> SEQ ID NO: 9
246 <211> LENGTH: 1083
247 <212> TYPE: DNA

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/580,906

DATE: 06/08/2006
TIME: 10:08:06

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

Use of <220> Feature (NEW RULES): *error explanation*
Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:5,6,7,10,11,12,13,14,15,16,17,18,19,20

VERIFICATION SUMMARY

DATE: 06/08/2006

PATENT APPLICATION: US/10/580,906

TIME: 10:08:06

Input Set : A:\Sequence Listing for RECEPTOR FUNCTION REGULATING

AGENT.txt

Output Set: N:\CRF4\06082006\J580906.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:26 M:283 W: Missing Blank Line separator, <400> field identifier
L:78 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <400> field identifier
L:153 M:283 W: Missing Blank Line separator, <400> field identifier
L:177 M:283 W: Missing Blank Line separator, <220> field identifier
L:178 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:5, <213>
ORGANISM:Artificial Sequence
L:178 M:283 W: Missing Blank Line separator, <400> field identifier
L:178 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:178
L:184 M:283 W: Missing Blank Line separator, <220> field identifier
L:185 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:6, <213>
ORGANISM:Artificial Sequence
L:185 M:283 W: Missing Blank Line separator, <400> field identifier
L:185 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:6,Line#:185
L:191 M:283 W: Missing Blank Line separator, <220> field identifier
L:192 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:7, <213>
ORGANISM:Artificial Sequence
L:192 M:283 W: Missing Blank Line separator, <400> field identifier
L:192 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:192
L:198 M:283 W: Missing Blank Line separator, <400> field identifier
L:249 M:283 W: Missing Blank Line separator, <400> field identifier
L:273 M:283 W: Missing Blank Line separator, <220> field identifier
L:274 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:10, <213>
ORGANISM:Artificial Sequence
L:274 M:283 W: Missing Blank Line separator, <400> field identifier
L:274 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10,Line#:274
L:280 M:283 W: Missing Blank Line separator, <220> field identifier
L:281 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:11, <213>
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L:281 M:283 W: Missing Blank Line separator, <400> field identifier
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L:287 M:283 W: Missing Blank Line separator, <220> field identifier
L:288 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:12, <213>
ORGANISM:Artificial Sequence
L:288 M:283 W: Missing Blank Line separator, <400> field identifier
L:288 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:12,Line#:288
L:294 M:283 W: Missing Blank Line separator, <220> field identifier
L:295 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:13, <213>
ORGANISM:Artificial Sequence
L:295 M:283 W: Missing Blank Line separator, <400> field identifier
L:295 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:295
L:301 M:283 W: Missing Blank Line separator, <220> field identifier
L:302 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:14, <213>
ORGANISM:Artificial Sequence
L:302 M:283 W: Missing Blank Line separator, <400> field identifier
L:302 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:14,Line#:302
L:308 M:283 W: Missing Blank Line separator, <220> field identifier
L:309 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:15, <213>
ORGANISM:Artificial Sequence
L:309 M:283 W: Missing Blank Line separator, <400> field identifier
L:309 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:309

L:315 M:283 W: Missing Blank Line separator, <220> field identifier
L:316 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:16, <213>
ORGANISM:Artificial Sequence
L:316 M:283 W: Missing Blank Line separator, <400> field identifier
L:316 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:16,Line#:316
L:322 M:283 W: Missing Blank Line separator, <220> field identifier